



IFW16

RAW SEQUENCE LISTING

DATE: 10/01/2004

PATENT APPLICATION: US/09/920,137B

TIME: 10:46:27

Input Set : A:\Cen 0250 SEQ LIST 07-09-01.txt

Output Set: N:\CRF4\10012004\I920137B.raw

5 <110> APPLICANT: Giles-Komar, Jill
 6 David Shealy
 7 David Knight
 8 Bernie Scallon
 9 George Heavner
 11 <120> TITLE OF INVENTION: ANTI-TNF ANTIBODIES, COMPOSITIONS, METHODS AND USES
 13 <130> FILE REFERENCE: CEN0250
 C--> 15 <140> CURRENT APPLICATION NUMBER: US/09/920,137B
 C--> 15 <141> CURRENT FILING DATE: 2001-08-01
 15 <160> NUMBER OF SEQ ID NOS: 15
 17 <170> SOFTWARE: PatentIn Ver 3.1

ERRORED SEQUENCES

19 <210> SEQ ID NO: 1
 20 <211> LENGTH: 5
 21 <212> TYPE: PRT
 22 <213> ORGANISM: Homo sapiens
 W--> 23 <400> SEQUENCE: 1
 25 Arg Tyr Thr Met His
 E--> 26 1 x 5 x
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 29 <211> LENGTH: 17
 30 <212> TYPE: PRT
 31 <213> ORGANISM: Homo sapiens
 W--> 32 <400> SEQUENCE: 2
 34 Val Ile Ser. Phe. Asp Gly Ser Asn Lys Tyr Tyr Val Asp Ser Val Lys
 E--> 35 1 x 5 x 10 10 15 x
 37 <210> SEQ ID NO: 3
 38 <211> LENGTH: 10
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 40 <213> ORGANISM: Homo sapiens
 W--> 41 <400> SEQUENCE: 3
 43 Glu Ala Arg Gly Ser Tyr Ala Phe Asp Ile
 E--> 44 1 x 5 x 10 10
 64 <210> SEQ ID NO: 6
 65 <211> LENGTH: 10
 66 <212> TYPE: PRT
 67 <213> ORGANISM: Homo sapiens
 W--> 68 <400> SEQUENCE: 6
 70 Gln Gln Arg Ser Asn Trp Pro Pro Phe Thr
 E--> 71 1 x 5 5 10 x

Does Not Comply
 Corrected Diskette Needed

(PS-1-3)

← Pls see item
 ↓ # 3 on error
 summary
 sheet.

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Input Set : A:\Cen 0250 SEQ LIST 07-09-01.txt

Output Set: N:\CRF4\10012004\I920137B.raw

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194 <212> TYPE: DNA
195 <213> ORGANISM: Homo sapiens
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Found 42

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VERIFICATION SUMMARY

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Input Set : A:\Cen 0250 SEQ LIST 07-09-01.txt

Output Set: N:\CRF4\10012004\I920137B.raw

L:15 M:270 C: Current Application Number differs, Replaced Current Application No
L:15 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:23 M:283 W: Missing Blank Line separator, <400> field identifier
L:26 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1 ✓
L:32 M:283 W: Missing Blank Line separator, <400> field identifier
L:35 M:252 E: No. of Seq. differs, <211> LENGTH:Input:17 Found:16 SEQ:2 ✓
L:41 M:283 W: Missing Blank Line separator, <400> field identifier ✓
L:44 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:3 ✓
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L:196 M:283 W: Missing Blank Line separator, <400> field identifier
L:198 M:254 E: No. of Bases conflict, LENGTH:Input:32 Counted:42 SEQ:13 ✓
L:198 M:252 E: No. of Seq. differs, <211> LENGTH:Input:32 Found:42 SEQ:13 ✓
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L:212 M:283 W: Missing Blank Line separator, <400> field identifier